

The Transcriptional Regulation of B Cell Lineage Commitment

Stephen L. Nutt* and Barbara L. Kee*

*Correspondence: nutt@wehi.edu.au (S.L.N.), bkee@bsd.uchicago.edu (B.L.K.)

DOI 10.1016/j.immuni.2007.07.006

(Immunity 27, 715–725; June 2007)

The authors regret that they inadvertently omitted citation of papers that were the first to propose an integrated model of early B cell development, a model that describes the interplay between the transcription factors controlling B cell development and cell-surface-receptor signaling pathways, from the Singh laboratory ([Medina et al., 2004](#); [Singh et al., 2005](#)). Figure 1 in their review was adapted from these papers and expanded to incorporate more recent findings.

Medina, K.L., Pongubala, J.M., Reddy, K.L., Lancki, D.W., Dekoter, R., Kieslinger, M., Grosschedl, R., and Singh, H. (2004). Assembling a gene regulatory network for specification of the B cell fate. *Dev. Cell* 7, 607–617.

Singh, H., Medina, K.L., and Pongubala, J.M. (2005). Contingent gene regulatory networks and B cell fate specification. *Proc. Natl. Acad. Sci. USA* 102, 4949–4953.